

* * * * * * * * * * * * * * * * STN Columbus * * * * * * * * * * * * * * * *

FILE 'HOME' ENTERED AT 15:16:59 ON 04 FEB 2010

=> fil .bec
COST IN U.S. DOLLARS
FULL ESTIMATED COST

| SINCE FILE ENTRY | TOTAL SESSION |
|------------------|---------------|
| 0.22 | 0.22 |

FILES 'MEDLINE, SCISEARCH, LIFESCI, BIOTECHDS, BIOSIS, EMBASE, HCAPLUS, NTIS,
ESBIOBASE, BIOTECHNO, WPIDS' ENTERED AT 15:17:29 ON 04 FEB 2010
ALL COPYRIGHTS AND RESTRICTIONS APPLY. SEE HELP USAGETERMS FOR DETAILS.

11 FILES IN THE FILE LIST

=> s (photoprotein# or aequorin or obelin or mnemiopsin or phiallidin or
mitrocomin or halistaurin or clytin)(15a) (muta? or variant#)

FILE 'MEDLINE'

| | |
|--------|---|
| 479 | PHOTOPROTEIN# |
| 1556 | AEQUORIN |
| 87 | OBELIN |
| 6 | MNEMIOPSIN |
| 0 | PHIALLIDIN |
| 3 | MITROCOMIN |
| 3 | HALISTAURIN |
| 6 | CLYTIN |
| 663898 | MUTA? |
| 152462 | VARIANT# |
| L1 | 48 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
) |

FILE 'SCISEARCH'

| | |
|--------|---|
| 503 | PHOTOPROTEIN# |
| 1510 | AEQUORIN |
| 96 | OBELIN |
| 6 | MNEMIOPSIN |
| 0 | PHIALLIDIN |
| 3 | MITROCOMIN |
| 1 | HALISTAURIN |
| 7 | CLYTIN |
| 661718 | MUTA? |
| 179116 | VARIANT# |
| L2 | 59 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
) |

FILE 'LIFESCI'

| | |
|--------|---|
| 210 | PHOTOPROTEIN# |
| 567 | AEQUORIN |
| 27 | OBELIN |
| 1 | MNEMIOPSIN |
| 0 | PHIALLIDIN |
| 2 | MITROCOMIN |
| 1 | HALISTAURIN |
| 5 | CLYTIN |
| 320581 | MUTA? |
| 61115 | VARIANT# |
| L3 | 25 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
) |

FILE 'BIOTECHDS'
77 PHOTOPROTEIN#
145 AEQUORIN
24 OBELIN
6 MNEMIOPSIN
0 PHIALLIDIN
3 MITROCOMIN
0 HALISTAURIN
12 CLYTIN
54258 MUTA?
19901 VARIANT#
L4 21 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

FILE 'BIOSIS'
593 PHOTOPROTEIN#
1818 AEQUORIN
119 OBELIN
5 MNEMIOPSIN
0 PHIALLIDIN
4 MITROCOMIN
2 HALISTAURIN
8 CLYTIN
730472 MUTA?
158947 VARIANT#
L5 64 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

FILE 'EMBASE'
532 PHOTOPROTEIN#
1356 AEQUORIN
56 OBELIN
4 MNEMIOPSIN
0 PHIALLIDIN
1 MITROCOMIN
1 HALISTAURIN
3 CLYTIN
568094 MUTA?
134196 VARIANT#
L6 35 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

FILE 'HCAPLUS'
764 PHOTOPROTEIN#
1661 AEQUORIN
147 OBELIN
12 MNEMIOPSIN
0 PHIALLIDIN
12 MITROCOMIN
3 HALISTAURIN
21 CLYTIN
682003 MUTA?
153947 VARIANT#
L7 94 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

FILE 'NTIS'

16 PHOTOPROTEIN#
21 AEQUORIN
2 OBELIN
0 MNEMIOPSIN
0 PHIALLIDIN
0 MITROCOMIN
0 HALISTAURIN
0 CLYTIN
11072 MUTA?
5176 VARIANT#
L8 3 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

FILE 'ESBIOBASE'
178 PHOTOPROTEIN#
604 AEQUORIN
33 OBELIN
0 MNEMIOPSIN
0 PHIALLIDIN
1 MITROCOMIN
0 HALISTAURIN
3 CLYTIN
365733 MUTA?
72121 VARIANT#
L9 42 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

FILE 'BIOTECHNO'
260 PHOTOPROTEIN#
416 AEQUORIN
27 OBELIN
0 MNEMIOPSIN
0 PHIALLIDIN
1 MITROCOMIN
1 HALISTAURIN
2 CLYTIN
242571 MUTA?
41198 VARIANT#
L10 20 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

FILE 'WPIDS'
122 PHOTOPROTEIN#
250 AEQUORIN
45 OBELIN
17 MNEMIOPSIN
0 PHIALLIDIN
13 MITROCOMIN
2 HALISTAURIN
22 CLYTIN
43447 MUTA?
39327 VARIANT#
L11 26 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDI
N OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#
)

TOTAL FOR ALL FILES
L12 437 (PHOTOPROTEIN# OR AEQUORIN OR OBELIN OR MNEMIOPSIN OR PHIALLIDIN
OR MITROCOMIN OR HALISTAURIN OR CLYTIN) (15A) (MUTA? OR VARIANT#)

```
=> s (phiallidin or clytin) and (muta? or variant#)
FILE 'MEDLINE'
    0 PHIALLIDIN
    6 CLYTIN
    663898 MUTA?
    152462 VARIANT#
L13          0 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'SCISEARCH'
    0 PHIALLIDIN
    7 CLYTIN
    661718 MUTA?
    179116 VARIANT#
L14          0 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'LIFESCI'
    0 PHIALLIDIN
    5 CLYTIN
    320581 MUTA?
    61115 VARIANT#
L15          0 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'BIOTECHDS'
    0 PHIALLIDIN
    12 CLYTIN
    54258 MUTA?
    19901 VARIANT#
L16          7 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'BIOSIS'
    0 PHIALLIDIN
    8 CLYTIN
    730472 MUTA?
    158947 VARIANT#
L17          0 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'EMBASE'
    0 PHIALLIDIN
    3 CLYTIN
    568094 MUTA?
    134196 VARIANT#
L18          0 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'HCAPLUS'
    0 PHIALLIDIN
    21 CLYTIN
    682003 MUTA?
    153947 VARIANT#
L19          5 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'NTIS'
    0 PHIALLIDIN
    0 CLYTIN
    11072 MUTA?
    5176 VARIANT#
L20          0 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'ESBIOBASE'
    0 PHIALLIDIN
    3 CLYTIN
```

365733 MUTA?
72121 VARIANT#
L21 2 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'BIOTECHNO'
0 PHIALLIDIN
2 CLYTIN
242571 MUTA?
41198 VARIANT#
L22 0 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

FILE 'WPIDS'
0 PHIALLIDIN
22 CLYTIN
43447 MUTA?
39327 VARIANT#
L23 10 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

TOTAL FOR ALL FILES
L24 24 (PHIALLIDIN OR CLYTIN) AND (MUTA? OR VARIANT#)

=> s (l12 or l24) not 2007-2010/pY

FILE 'MEDLINE'
2199154 2007-2010/PY
L25 35 (L1 OR L13) NOT 2007-2010/PY

FILE 'SCISEARCH'
4083879 2007-2010/PY
(20070000-20109999/PY)
L26 44 (L2 OR L14) NOT 2007-2010/PY

FILE 'LIFESCI'
696093 2007-2010/PY
L27 18 (L3 OR L15) NOT 2007-2010/PY

FILE 'BIOTECHDS'
50427 2007-2010/PY
L28 18 (L4 OR L16) NOT 2007-2010/PY

FILE 'BIOSIS'
1803119 2007-2010/PY
L29 47 (L5 OR L17) NOT 2007-2010/PY

FILE 'EMBASE'
1832835 2007-2010/PY
L30 26 (L6 OR L18) NOT 2007-2010/PY

FILE 'HCAPLUS'
4894016 2007-2010/PY
L31 62 (L7 OR L19) NOT 2007-2010/PY

FILE 'NTIS'
45930 2007-2010/PY
L32 3 (L8 OR L20) NOT 2007-2010/PY

FILE 'ESBIOBASE'
1072620 2007-2010/PY
L33 32 (L9 OR L21) NOT 2007-2010/PY

FILE 'BIOTECHNO'
0 2007-2010/PY
L34 20 (L10 OR L22) NOT 2007-2010/PY

FILE 'WPIDS'
4739359 2007-2010/PY
L35 9 (L11 OR L23) NOT 2007-2010/PY

TOTAL FOR ALL FILES
L36 314 (L12 OR L24) NOT 2007-2010/PY

=> dup rem 136
PROCESSING COMPLETED FOR L36
L37 105 DUP REM L36 (209 DUPLICATES REMOVED)

=> d tot

L37 ANSWER 1 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI New photoprotein aequorin Y89F, its functional equivalents and
corresponding nucleic acids, useful as labels or reporters, especially in
pharmacological research and diagnostic applications;
DNA and RNA vector-mediated gene transfer and expression in host cell
for use as a diagnostic, in pharmacological and pharmaceutical
industries, for calcium concentration detection, drug screening and
high throughput screening
AU GOLZ S; VYSOTSKI E; MARKOVA S; STEPANYUK G; BURAKOVA L; FRANK L
AN 2006-08904 BIOTECHDS
PI WO 2006010454 2 Feb 2006

L37 ANSWER 2 OF 105 HCPLUS COPYRIGHT 2010 ACS on STN DUPLICATE 2
TI Extending light-emitting time of calcium-binding photoprotein solution.
SO Brit. UK Pat. Appl., 45pp.
CODEN: BAXXDU
IN Inouye, Satoshi; Sasaki, Satoko
AN 2006:1145319 HCPLUS
DN 145:467722
PATENT NO. KIND DATE APPLICATION NO. DATE
----- ----- ----- -----
PI GB 2425535 A 20061101 GB 2006-8514 20060428
JP 2006308501 A 20061109 JP 2005-133743 20050428
US 20060246534 A1 20061102 US 2006-411715 20060426

L37 ANSWER 3 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI Photoprotein useful as an intracellular calcium indicator, is
obtained by mutagenesis of clytin, is able to bind
coelenterazine and calcium, and displays enhanced bioluminescence;
recombinant photoprotein for intracellular calcium indicator,
cell-based high throughput screening assay, intracellular calcium
concentration modulating compound screening and diagnosis composition
AU MASTROIANNI N; CAINARCA S; CORAZZA S
AN 2006-24828 BIOTECHDS
PI WO 2006094805 14 Sep 2006

L37 ANSWER 4 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI Novel cell comprising endogenous promiscuous G-proteins and exogenous
nucleic acid encoding G-protein coupled receptor (GPCR), useful for
identifying agent that modulates activity of GPCR;
a recombinant G-protein coupled receptor expressed in a Chinese
hamster ovary cell useful for the identification of an agonist or
antagonist
AU HSU M
AN 2006-13695 BIOTECHDS
PI WO 2006050214 11 May 2006

L37 ANSWER 5 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN

TI New nucleic acid encoding variant of aequorin, useful
e.g. as reporter gene and as dye, has much longer luminescent lifetime
than the parent protein, also new encoded proteins;
recombinant photoprotein production via plasmid expression in host
cell for use in marker and pollutant quantification
AN 2007-01755 BIOTECHDS
PI DE 102005022146 23 Nov 2006

L37 ANSWER 6 OF 105 HCPLUS COPYRIGHT 2010 ACS on STN
TI Genetically engineered luminescent proteins in biosensing
SO Proceedings of SPIE-The International Society for Optical Engineering
(2006), 6098, 60980H/1-60980H/9
CODEN: PSISDG; ISSN: 0277-786X
AU Rowe, Laura; Ensor, Mark; Scott, Daniel; Deo, Sapna; Daunert, Sylvia
AN 2006:293863 HCPLUS
DN 145:287941

L37 ANSWER 7 OF 105 MEDLINE on STN DUPLICATE 3
TI Calcium dependence of aequorin bioluminescence dissected by
random mutagenesis.
SO Proceedings of the National Academy of Sciences of the United States of
America, (2006 Jun 20) Vol. 103, No. 25, pp. 9500-5. Electronic
Publication: 2006-06-12.
Journal code: 7505876. ISSN: 0027-8424. L-ISSN: 0027-8424.
Report No.: NLM-PMC1480436.
AU Tricoire Ludovic; Tsuzuki Keisuke; Courjean Olivier; Gibelin Nathalie;
Bourout Gaelle; Rossier Jean; Lambolez Bertrand
AN 2006373143 MEDLINE

L37 ANSWER 8 OF 105 MEDLINE on STN DUPLICATE 4
TI Crystal structure of obelin after Ca²⁺-triggered bioluminescence suggests
neutral coelenteramide as the primary excited state.
SO Proceedings of the National Academy of Sciences of the United States of
America, (2006 Feb 21) Vol. 103, No. 8, pp. 2570-5. Electronic
Publication: 2006-02-08.
Journal code: 7505876. ISSN: 0027-8424. L-ISSN: 0027-8424.
Report No.: NLM-PMC1413834.
AU Liu Zhi-Jie; Stepanyuk Galina A; Vysotski Eugene S; Lee John; Markova
Svetlana V; Malikova Natalia P; Wang Bi-Cheng
AN 2006245287 MEDLINE

L37 ANSWER 9 OF 105 SCISEARCH COPYRIGHT (c) 2010 The Thomson Corporation on
STN DUPLICATE 5
TI Presenilin mutations linked to familial Alzheimer's disease reduce
endoplasmic reticulum and Golgi apparatus calcium levels
SO CELL CALCIUM, (JUN 2006) Vol. 39, No. 6, pp. 539-550.
ISSN: 0143-4160.
AU Pizzo P (Reprint); Zatti G; Burgo A; Giacomello M; Barbiero L; Ghidoni R;
Sinigaglia G; Florean C; Bagnoli S; Binetti G; Sorbi S; Fasolato C
AN 2006:638192 SCISEARCH

L37 ANSWER 10 OF 105 SCISEARCH COPYRIGHT (c) 2010 The Thomson Corporation
on STN DUPLICATE 6
TI Calcium dependence of aequorin bioluminescence dissected by
random mutagenesis
SO LUMINESCENCE, (SEP-OCT 2006) Vol. 21, No. 5, pp. 280-281.
ISSN: 1522-7235.
AU Lambolez, B. (Reprint); Tricoire, L.; Tsuzuki, K.
AN 2007:63333 SCISEARCH

L37 ANSWER 11 OF 105 MEDLINE on STN DUPLICATE 7
TI Photoprotein aequorin as a novel reporter for SNP

- genotyping by primer extension-application to the variants of mannose-binding lectin gene.
- SO Human mutation, (2006 Mar) Vol. 27, No. 3, pp. 279-85.
Journal code: 9215429. E-ISSN: 1098-1004. L-ISSN: 1059-7794.
- AU Zerefos Panayotis G; Ioannou Penelope C; Traeger-Synodinos Joanne; Dimissianos Gerasimos; Kanavakis Emmanuel; Christopoulos Theodore K
- AN 2006090326 MEDLINE
- L37 ANSWER 12 OF 105 SCISEARCH COPYRIGHT (c) 2010 The Thomson Corporation on STN
DUPLICATE 8
- TI Obelin mutants with altered colour of light emission as labels for dual-wavelength immunoassay
- SO LUMINESCENCE, (SEP-OCT 2006) Vol. 21, No. 5, pp. 271-271.
ISSN: 1522-7235.
- AU Borisova, V. V. (Reprint); Frank, L. A.; Malikova, N. P.; Stepanyuk, G. A.; Markova, S. V.; Lee, J.; Vysotski, E. S.
- AN 2007:63300 SCISEARCH
- L37 ANSWER 13 OF 105 HCAPLUS COPYRIGHT 2010 ACS on STN
- TI Luminescent proteins in binding assays
- SO Photoproteins in Bioanalysis (2006), 155-178. Editor(s): Daunert, Sylvia; Deo, Sapna K. Publisher: Wiley-VCH Verlag GmbH & Co. KGaA, Weinheim, Germany.
CODEN: 69IPFP; ISBN: 978-3-527-31016-6
- AU Roda, Aldo; Guardigli, Massimo; Michelini, Elisa; Mirasoli, Mara; Pasini, Patrizia
- AN 2006:1159037 HCAPLUS
- DN 146:223628
- L37 ANSWER 14 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
- TI Monitoring the expression level of a gene comprises transforming a cell expressing a regulatory biomolecule with a nucleic acid molecule encoding an interaction partner of the biomolecule;
gene expression level monitoring via vector expression in host cell
- AU HILLEN W; BERENS C; KLOTZSCHE M
- AN 2005-26651 BIOTECHDS
- PI EP 1580273 28 Sep 2005
- L37 ANSWER 15 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
- TI Monitoring the expression level of a gene in a host cell by modulating a regulatory biomolecule activity by transforming a cell expressing a regulatory biomolecule with a nucleic acid and assessing the expression level of the gene;
for use in gene expression monitoring
- AU HILLEN W; KLOTZSCHE M; BERENS C
- AN 2005-28936 BIOTECHDS
- PI WO 2005093075 6 Oct 2005
- L37 ANSWER 16 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
- TI Detecting dynamics of calcium ion in biological system, optically, involves monitoring photons emitted by recombinant calcium-sensitive polypeptide having chemiluminescent protein linked to fluorescent protein, present in system;
transgenic animal model construction production via plasmid expression in host cell for use in disease diagnosis and calcium ion detection
- AU BRULET P; ROGERS K; PICAUD S
- AN 2005-25434 BIOTECHDS
- PI WO 2005078445 25 Aug 2005
- L37 ANSWER 17 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
- TI New isolated nucleic acid encoding aequorin or obelin mutant protein capable of binding coelenterazine and molecular

- oxygen, and emitting light, useful for multianalyte microanalysis, and for identifying inhibitors of HIV-1 protease;
mutant protein molecule isolation for use in microanalysis and virus enzyme-inhibitor identification
- AU DAUNERT S; DEO S K; DIKICI E; ROWE L
AN 2005-29507 BIOTECHDS
PI US 20050214776 29 Sep 2005
- L37 ANSWER 18 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI New nucleic acid encoding photoproteins from Clytia gregaria, useful as marker and reporter genes, particularly in screening for pharmaceuticals;; recombinant protein production via plasmid expression in host cell for use in marker and reporter gene
- AU GOLZ S; MARKOVA S; BURAKOVA L; FRANK L; VYSOTSKI E
AN 2005-15281 BIOTECHDS
PI DE 10342670 21 Apr 2005
- L37 ANSWER 19 OF 105 MEDLINE on STN DUPLICATE 10
TI Thermostable mutants of the photoprotein aequorin obtained by in vitro evolution.
SO The Journal of biological chemistry, (2005 Oct 7) Vol. 280, No. 40, pp. 34324-31. Electronic Publication: 2005-06-22. Journal code: 2985121R. ISSN: 0021-9258. L-ISSN: 0021-9258.
- AU Tsuzuki Keisuke; Tricoire Ludovic; Courjean Olivier; Gibelin Nathalie; Rossier Jean; Lambolez Bertrand
AN 2005531200 MEDLINE
- L37 ANSWER 20 OF 105 MEDLINE on STN DUPLICATE 11
TI Transient receptor potential-like channels are essential for calcium signaling and fluid transport in a Drosophila epithelium.
SO Genetics, (2005 Mar) Vol. 169, No. 3, pp. 1541-52. Electronic Publication: 2005-02-03. Journal code: 0374636. ISSN: 0016-6731. L-ISSN: 0016-6731. Report No.: NLM-PMC1449567.
- AU MacPherson Matthew R; Pollock Valerie P; Kean Laura; Southall Tony D; Giannakou Maria E; Broderick Kate E; Dow Julian A T; Hardie Roger C; Davies Shireen A
AN 2005162895 MEDLINE
- L37 ANSWER 21 OF 105 MEDLINE on STN DUPLICATE 12
TI Bioluminescence resonance energy transfer from aequorin to a fluorophore: an artificial jellyfish for applications in multianalyte detection.
SO Analytical and bioanalytical chemistry, (2005 Apr) Vol. 381, No. 7, pp. 1387-94. Electronic Publication: 2005-02-25. Journal code: 101134327. ISSN: 1618-2642.
- AU Deo Sapna K; Mirasoli Mara; Daunert Sylvia
AN 2005186632 MEDLINE
- L37 ANSWER 22 OF 105 LIFESCI COPYRIGHT 2010 CSA on STN DUPLICATE 13
TI Interchange of aequorin and obelin bioluminescence color is determined by substitution of one active site residue of each photoprotein
SO FEBS Letters [FEBS Lett.], (20050200) vol. 579, no. 5, pp. 1008-1014. ISSN: 0014-5793.
- AU Stepanyuk, Galina A; Golz, Stefan; Markova, Svetlana V; Frank, Ludmila A; Lee, John; Vysotski, Eugene S
AN 2007:220464 LIFESCI
- L37 ANSWER 23 OF 105 MEDLINE on STN DUPLICATE 14
TI Motilin and erythromycin-A share a common binding site in the third transmembrane segment of the motilin receptor.
SO Biochemical pharmacology, (2005 Sep 15) Vol. 70, No. 6, pp. 879-87.

Journal code: 0101032. ISSN: 0006-2952. L-ISSN: 0006-2952.
AU Xu Luo; Depoortere Inge; Vertogenen Pascale; Waelbroeck Magali; Robberecht Patrick; Peeters Theo L
AN 2005431805 MEDLINE

L37 ANSWER 24 OF 105 Elsevier Biobase COPYRIGHT 2010 Elsevier Science B.V. on STN
AN 2005115792 ESBIOBASE
TI Suppression of Pdx-1 perturbs proinsulin processing, insulin secretion and GLP-1 signalling in INS-1 cells
AU Wang, H.; Iezzi, M.; Theander, S.; Antinozzi, P.A.; Gauthier, B.R.; Wollheim, C.B.; Halban, P.A.
CS Wang, H.; Iezzi, M.; Theander, S.; Antinozzi, P.A.; Gauthier, B.R.; Wollheim, C.B. (Dept. of Cell Physiol. and Metab., University Medical Center, 1211 Geneva 4 (CH)); Halban, P.A. (Dept. of Med. Genet. and Development, University Medical Center, Geneva (CH))
EMAIL: Haiyan.Wang@medicine.unige.ch
SO Diabetologia (Apr 2005) Volume 48, Number 4, pp. 720-731, 71 refs.
CODEN: DBTGAI ISSN: 0012-186X
DOI: 10.1007/s00125-005-1692-8
CY Germany
DT Journal; Article
LA English
SL English
ED Entered STN: 3 Feb 2009
Last updated on STN: 3 Feb 2009

L37 ANSWER 25 OF 105 MEDLINE on STN DUPLICATE 15
TI Effect of inactivating mutations on phosphorylation and internalization of the human VPAC2 receptor.
SO Journal of molecular endocrinology, (2005 Apr) Vol. 34, No. 2, pp. 405-14.
Journal code: 8902617. ISSN: 0952-5041. L-ISSN: 0952-5041.
AU Langer Ingrid; Langlet Christelle; Robberecht Patrick
AN 2005187883 MEDLINE

L37 ANSWER 26 OF 105 HCAPLUS COPYRIGHT 2010 ACS on STN
TI Spectral tuning of the bioluminescent photoprotein Aequorin
SO Abstracts of Papers, 229th ACS National Meeting, San Diego, CA, United States, March 13-17, 2005 (2005), ANYL-179 Publisher: American Chemical Society, Washington, D. C.
CODEN: 69GQMP
AU Rowe, Laura; Dikici, Emre; Logue, Courtney; Scott, Daniel; Deo, Sapna; Daunert, Sylvia
AN 2005:185996 HCAPLUS

L37 ANSWER 27 OF 105 SCISEARCH COPYRIGHT (c) 2010 The Thomson Corporation on STN DUPLICATE 16
TI Pharmacological investigation of the Arg(344)His variant of the human 5-HT3A receptor by radioligand-binding and aequorin-based calcium-influx measurement
SO NAUNYN-SCHMIEDEBERGS ARCHIVES OF PHARMACOLOGY, (FEB 2005) Vol. 371, Supp. [1], pp. R30-R30. MA 120.
ISSN: 0028-1298.
AU Combrink S (Reprint); Kostanian A; Barann M; Bonisch H; Gothert M; Bruss M
AN 2005:751948 SCISEARCH

L37 ANSWER 28 OF 105 LIFESCI COPYRIGHT 2010 CSA on STN
TI Transient Receptor Potential-Like Channels Are Essential for Calcium Signaling and Fluid Transport in a Drosophila Epithelium
SO Genetics, (20050300) vol. 169, no. 3, [np].
ISSN: 0016-6731.
AU MacPherson, Matthew R.; Pollock, Valerie P.; Kean, Laura; Southall, Tony

- D.; Giannakou, Maria E.; Broderick, Kate E.; Dow, Julian A. T.; Hardie, Roger C.; Davies, Shireen A.
- AN 2007:146194 LIFESCI
- L37 ANSWER 29 OF 105 MEDLINE on STN DUPLICATE 17
TI Self-reporting Arabidopsis expressing pH and [Ca²⁺] indicators unveil ion dynamics in the cytoplasm and in the apoplast under abiotic stress.
SO Plant physiology, (2004 Mar) Vol. 134, No. 3, pp. 898-908.
Journal code: 0401224. ISSN: 0032-0889. L-ISSN: 0032-0889.
Report No.: NLM-PMC389913.
- AU Gao Dongjie; Knight Marc R; Trewavas Anthony J; Sattelmacher Burkhard;
Plieth Christoph
AN 2004129591 MEDLINE
- L37 ANSWER 30 OF 105 MEDLINE on STN DUPLICATE 18
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AN 2005060482 MEDLINE
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TI Analysis of the Ca²⁺ response of mycelial fungi to external effects by the recombinant aequorin method
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El'-Registan G I
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Rose John; Wang Bi-Cheng
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TI New mutated photoproteins derived from jellyfish photoproteins have increased thermostability or increased luminescence time and are useful as bioluminescent markers, e.g., to detect pathogens;
vector-mediated gene transfer and expression in Escherichia coli,
HEK-293 or CHO cell for HIV virus infection diagnosis

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ROSSIER J
AN 2003-13579 BIOTECHDS
PI FR 2827292 17 Jan 2003

L37 ANSWER 35 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI Novel recombinant calcium-binding photoprotein useful for producing
conjugates which in turn is useful as marker in immunoassay;
vector-mediated gene transfer and expression in host cell for
recombinant protein production and immunoassay marker

AU INOUYE S
AN 2004-03144 BIOTECHDS
PI US 20030212259 13 Nov 2003

L37 ANSWER 36 OF 105 HCAPLUS COPYRIGHT 2010 ACS on STN
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bioluminescence for high-throughput screening of calcium ion flux in cell
SO PCT Int. Appl., 46 pp.
CODEN: PIXXD2

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DN 139:287962

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CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR,
LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH,
PL, PT, RO, RU, SD, SE, SG, SK, SL, TJ, TM, TN, TR, TT, TZ, UA,
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RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AM, AZ, BY,
KG, KZ, MD, RU, TJ, TM, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES,
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L37 ANSWER 37 OF 105 HCAPLUS COPYRIGHT 2010 ACS on STN
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luminescence lifetime, their production with recombinant cells, and their
uses in bioassays

SO PCT Int. Appl., 184 pp.
CODEN: PIXXD2

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Eric; Courjean, Olivier Arsene; Tsuzuki, Keisuke; Rossier, Jean

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DN 138:119118

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RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AM, AZ, BY,
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 SO Journal of Muscle Research and Cell Motility (2003), Volume Date 2002, 23(7-8), 853-865
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vector-mediated gene transfer and expression in yeast host cell for recombinant protein production
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vector-mediated gene transfer and expression in host cell for drug screening
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vector-mediated reporter gene transfer, expression in host cell and antibody for recombinant protein production
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PI EP 1156103 21 Nov 2001

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JP 2003525596 W 20030902 (200358) JA 80
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L37 ANSWER 88 OF 105 HCAPLUS COPYRIGHT 2010 ACS on STN

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prepared by site-specific mutagenesis process; expression in Escherichia coli
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L37 ANSWER 104 OF 105 SCISEARCH COPYRIGHT (c) 2010 The Thomson Corporation on STN DUPLICATE 53
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L37 ANSWER 105 OF 105 BIOSIS COPYRIGHT (c) 2010 The Thomson Corporation on STN

TI PCR-based *in vitro* mutagenesis of the photoprotein aequorin.

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=> d ab 6,17,19,26,32,33,35,36,43,87,88,95,105

L37 ANSWER 6 OF 105 HCPLUS COPYRIGHT 2010 ACS on STN

AB Luminescent proteins originally isolated from marine or terrestrial organisms have played a key role in the development of several biosensing systems. These proteins have been used in a variety of applications including, immunoassays, binding assays, cell-based sensing, high throughput screening, optical imaging, etc. Among the luminescent proteins isolated, the bioluminescent protein aequorin has been one of the proteins at the forefront in terms of its use in a vast number of biosensing systems. In our laboratory, we have employed aequorin as a label in the development of highly sensitive assays through chemical and genetic modifications from single step anal. of physiol. important mols. in biol. fluids. An important aspect of optimizing these assays for clin. use involves understanding the stability of the various aequorin variants that are available. To this end we have designed several stability studies involving three important aequorin mutants, Mutant S, Mutant 5, and Mutant 53. The cysteine free aequorin, Mutant S, has been the most ubiquitously used aequorin variant in our laboratory because of its increased stability and activity as compared to native aequorin. Mutant 5 and Mutant 53 contain a single cysteine residue at position 5 and 53 in the protein, resp. Because of the presence of a single cysteine residue, Mutant 5 and Mutant 53 both can be site-specifically conjugated. This site specific conjugation capability gives Mutant 5 and Mutant 53 an advantage over native aequorin when developing assays. Addnl. studies optimizing the expression, purification, and charging of aequorin Mutant S were also performed. A thorough understanding of the efficient expression, purification, and storage of these aequorin mutants will allow for the more practical utilization of these mutants in the development of future biosensing systems.

L37 ANSWER 17 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN

AB DERWENT ABSTRACT:

NOVELTY - An isolated nucleic acid (N1) hybridizing with a fully defined 861 or 662 base pairs (SEQ ID NO. 3 or 5) sequence given in the specification, under stringent conditions, and encoding a protein capable of binding coelenterazine and molecular oxygen, and emitting light, is new.

DETAILED DESCRIPTION - An isolated nucleic acid (N1) is chosen from:
(a) a nucleic acid hybridizing with a fully defined 861 base pairs (SEQ ID NO. 3) sequence given in the specification, under stringent conditions, and encoding a protein capable of binding coelenterazine and molecular oxygen, and emitting light, where (i) the protein has an

isoleucine residue in first position corresponding to position 132 of fully defined 189 amino acid (SEQ ID Number 4) sequence given in the specification; (ii) a non-natural amino acid is incorporated into a position corresponding to 132 of SEQ ID Number 4, during translation of the protein; (iii) the protein has a cysteine residue in a first position corresponding to positions 65, 66, 69, 70, 74 or 76 of SEQ ID Number 4; (iv) the protein has a phenylalanine residue in a first position corresponding to position 132 or 82 of SEQ ID Number 4; (v) the protein has an tyrosine residue in a first position corresponding to position 86 or 16 of SEQ ID Number 4; or (vi) the protein has a tryptophan residue in a first position corresponding to position 82 of SEQ ID Number 4; and (b) a nucleic acid hybridising with a fully defined 662 base pairs (SEQ ID NO. 5) sequence given in the specification, under stringent conditions, and encoding a protein capable of binding coelenterazine and molecular oxygen, and emitting light, where the protein has a serine residue in a first position corresponding to positions 51, 67, or 151, and a serine residue in a second position corresponding to position 75 of a fully defined 165 amino acid (SEQ ID NO. 6) sequence given in the specification.

INDEPENDENT CLAIMS are also included for: (1) a kit (K1) comprising the protein encoded by (N1), and a coelenterazine chosen from CTZ i, ip, h, hcp, cp, fcp, f, n and native coelenterazine; (2) an aequorin mutant protein encoded by (N1), where the protein is conjugated to a fluorophore; and (3) identifying (M1) inhibitors of bond-breaking enzymes, preferably HIV-1 protease, comprising: (a) immobilizing a fusion protein encoded by a fusion protein nucleic acid comprising (N1) being operably linked to second nucleic acid encoding a bond-breaking enzyme recognition site, in a first locus and a second locus; (b) contacting the fusion protein with a candidate compound in the presence of the bond-breaking enzyme in first locus; (c) contacting the fusion protein with the bond-breaking enzyme in the second locus; and (d) determining whether there is an increase in the intensity of light emission at the first locus relative to light emission in the second locus.

BIOTECHNOLOGY - Preferred Nucleic Acid: In (N1), the non-natural amino acid is fluorotyrosine or fluorotryptophan, preferably 5-fluoro-1-tryptophan or 3-fluoro-1-tyrosine. Preferred Protein: In the aequorin mutant protein, the fluorophore is N-(((2-iodoacetoxy)ethyl)-N-methyl)amino-7-nitrobenz-2-oxa-1,3-diazole (IANBD) ester. Preferred Method: In (M1), the fusion protein comprises a non-natural amino acid. The recognition site is Ser-Glu-Asn-Tyr-Pro-Ile-Val (SEQ ID Number 5).

ACTIVITY - Anti-HIV.

MECHANISM OF ACTION - HIV-1 protease inhibitor; HIV protease inhibitor.

USE - A protein encoded by (N1) useful in identifying inhibitors of bond-breaking enzymes, preferably HIV-1 protease (claimed). HIV-1 protease inhibitors may be used for treating HIV-1 infections. The protein is useful in multianalyte microanalysis, in development of different competitive and non-competitive assays for detecting physiologically important molecules such as peptides, drugs, etc.

ADVANTAGE - The protein encoded by (N1) is capable of emitting light at predictably shifted wavelength when used with various coelenterazine variants, and enables high-throughput screening of biopharmaceuticals.

EXAMPLE - Microorganisms such as *Escherichia coli* containing a plasmid having genetic information for aequorin were grown in a minimal media supplemented with essential amino acids and vitamins not having amino acid of the analogue that would be incorporated. After the cells were grown to a certain optical density the analogue and the inducer, isopropyl-beta-D-thiogalactopyranoside (IPTG), was added in order to express aequorin. As the protein expression has taken place, the amino acid residues were replaced by the analogue present in the medium. The percent incorporation depends largely on the type and nature of amino acid analogue used and can range from 10-75%. The cells were then

harvested by centrifuging and lysed by sonication. The cell debris was separated from the supernatant, and the supernatant was then incubated with different coelenterazine analogues and the emission spectra were noted. Results showed that an emission wavelength of 511 nm was observed, when the supernatant containing protein was incubated with coelenterazine. Thus, the supernatant containing aequorin mutants having non-natural amino acids was obtained.(35 pages)

- L37 ANSWER 19 OF 105 MEDLINE on STN DUPLICATE 10
- AB Aequorin is a photoprotein that emits light upon binding calcium. Aequorin mutants showing increased intensity or slow decay of bioluminescence were isolated by in vitro evolution combining DNA shuffling and functional screening in bacteria. Luminescence decay mutants were isolated at the first round of screening and carried mutations located in EF-hand calcium binding sites or their vicinity. During in vitro evolution, the luminescence intensity of the population of mutants increased with the frequency of effective mutations whereas the frequency of other amino acid substitutions remained roughly stable. Luminescence intensity mutations neighbored the His-16 or His-169 coelenterazine binding residues or were located in the first EF-hand. None of the selected mutants exhibited an increase in photon yield when examined in a cell-free assay. However, we observed that two mutants, Q168R and L170I, exhibited an increase of the photoprotein lifetime at 37 degrees C that may underlie their high luminescence intensity in bacteria. Further analysis of Q168R and L170I mutations showed that they increased aequorin thermostability. Conversely, examination of luminescence decay mutants revealed that the F149S substitution decreased aequorin thermostability. Finally, screening of a library of random Gln-168 and Leu-170 mutants confirmed the involvement of both positions in thermostability and indicated that optimal thermostability was conferred by Q168R and L170I mutations selected through in vitro evolution. Our results suggest that Phe-149 and Gln-168 residues participate in stabilization of the coelenterazine peroxide and the triggering of photon emission by linking the third EF-hand to Trp-129 and His-169 coelenterazine binding residues.

- L37 ANSWER 26 OF 105 HCPLUS COPYRIGHT 2010 ACS on STN
- AB Aequorin is a photoprotein whose calcium controlled bioluminescent light emission is used as a label in assays and for the determination of calcium concns. in vivo. Aequorin contains a coelenterazine chromophore which emits bioluminescence of a characteristic wavelength (465 nm) following relaxation from its oxidized state. Variants of the photoprotein aequorin, with shifted emission wavelengths, were prepared in order to enhance the applications of aequorin in bioanal. Here we show that combining four rationally designed aequorin mutants with various chromophore analogs results in significant spectral and half life shifts. Aequorin variants with different emission maxima and half lifes should allow aequorin to be employed in a variety of multi-analyte detection applications.

- L37 ANSWER 32 OF 105 MEDLINE on STN DUPLICATE 20
- AB Ca(2+)-regulated photoproteins belong to the EF-hand Ca(2+)-binding protein family. The addition of calcium ions initiates bright blue bioluminescence of the photoproteins, a result of the oxidative breakdown of coelenterazine peroxide to coelenteramide. Crystals of the Ca(2+)-discharged W92F mutant of obelin from Obelia longissima have been grown, representing the first crystallization of a photoprotein after the Ca(2+)-triggered bioluminescence. A green fluorescence observed from the crystals clearly demonstrates that

coelenteramide, the bioluminescence product of coelenterazine peroxide, is bound within the protein. The diffraction pattern exhibits tetragonal Laue symmetry. Systematic absences indicate that the space group is either P4(3)2(1)2 or P4(1)2(1)2. The unit-cell parameters are $a = b = 53.4$, $c = 144.0$ Å. The crystals diffract to 1.9 Å resolution.

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L37 ANSWER 35 OF 105 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
AB DERWENT ABSTRACT:

NOVELTY - A recombinant calcium-binding photoprotein (I) comprising a wild-type or mutant apoprotein having one cysteine residue introduced within the fourth amino acid residue from the amino terminus of the wild-type apoprotein, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for the following: (1) a DNA (II) encoding (I) comprising one cysteine residue introduced within the fourth amino acid residue from the amino terminus of the amino acid sequence of wild-type apoprotein having a full defined sequence (S1) of 189 amino acids as given in the specification or its mutant apoprotein; (2) an expression vector (III) comprising (II); (3) a transformed host cell (IV) comprising (III); (4) a conjugate (V) in which a ligand specific for a substance to be detected binds to (I) comprising a wild-type or mutant apoprotein having one cysteine residue introduced with the fourth amino acid residue from the amino terminus of the wild-type apoprotein, through the cysteine residue, in a ratio of 1:1; (5) producing (M1) (V), involves producing by genetically engineering an apoprotein having one cysteine residue introduced within the fourth amino acid residue from the amino terminus of (S1) or its mutant apoprotein, treating the apoprotein with coelenterazine in the presence of molecular oxygen to yield (I), and binding the photoprotein through the introduced cysteine residue to a ligand specific for a substance to be detected, in a ratio of 1:1; and (6) a kit (VI) for carrying out measuring a substance specific for a ligand, featuring by comprising (V).

BIOTECHNOLOGY - Preferred Photoprotein: (I) is chosen from aequorin, obelin, clytin, mitrocomin, mnemiopsin and berovin. (I) is present as luminescent substrate coelenterazine or its analog which exhibits the luminescence activity. (I) further comprises an apoprotein having a fully defined sequence (S2) of 192 amino acids as given in the specification, or an apoprotein having (S2) modified by a deletion, substitution or addition of 1 to 5 amino acids such that the sixth cysteine is conserved and the luminescence activity is maintained.

Preferred Conjugate: In (V), the ligand specific for a substance to be detected is biotin, avidin, streptavidin, an enzyme, a substrate, an antibody, an antigen, nucleic acid, a polysaccharide, a receptor or a compound capable of binding to any of these. The ligand specific for a substance to be detected binds to (I) comprising an apoprotein having one cysteine residue introduced within the fourth amino acid residue from the amino terminus of (S1) or its mutant apoprotein, through the introduced cysteine residue, in a ratio of 1:1. The ligand specific for a substance to be detected binds to (I) comprising an apoprotein having (S2), or an apoprotein having an amino acid sequence in which (S2) is modified by deletion, substitution or addition of 1 to 5 amino acids such that the sixth cysteine is conserved and luminescence activity is maintained, through the 6th cysteine, in a ratio of 1:1. The ligand specific for a substance to be detected is biotin.

Preferred Method: (M1) further involves producing by genetic engineering an apoprotein having (S2) or an apoprotein having modified (S2) in which the amino acid sequence is modified by a deletion, substitution or addition of 1 to 5 amino acids such that the sixth cysteine is conserved and luminescence activity is maintained, treating the apoprotein with coelenterazine in

the presence of oxygen to yield (I), and binding the photoprotein through the sixth cysteine to a ligand specific for a substance to be detected, in a ratio of 1:1.

USE - (V) is useful for measuring a substance specific for a ligand (claimed). (V) is useful as a marker for immunoassay.

EXAMPLE - From expression vector piP-HE with mutant apoaequorin (having an Ala-Asn-Ser sequence instead of valine at the N-terminus of the wild-type apoaequorin) the EcoRI restriction enzyme site (near the N-terminus of the apoaequorin gene) was deleted by PCR to construct piP-HEDELTAE. piP-HELE plasmid and PCR primers such as Cys4-Aq (5'GGCAAGCTTGTACTAGTGACTTCGACAACCCAAGATGG3') and 630EcoRI-AQ(5'GCC-GAA-TTC-ATC-AGT-GTT-TTA-TTC-AAA3') were used for PCR amplification of the target fragment with a geneAmp PCR reagent kit, followed by isolation of the fragment with a purification kit and digestion with restriction enzymes HindIII and EcoRI to obtain a HindIII-EcoRI fragment with cysteine at the sixth position from the N-terminus of the mutant apoaequorin gene (fourth position from the N-terminus of the wild-type apoaequorin gene). Separately, plasmid piP-HEDELTAE was digested with restriction enzymes HindIII and EcoRI, and the vector end containing the promoter and OmpA signal peptide was isolated. This was linked with the HindIII-EcoRI fragment and the obtained plasmid was used to transform E.coli JM83. Plasmid piP-HE-Cys4 expressing the mutant apoaequorin with cysteine inserted at the sixth position from the N-terminus was isolated from the transformants. The base sequence was determined and confirmed to be the cysteine-inserted apoaequorin (Cys4-apoaequorin). The amino acid sequence of Cys4-apoaequorin had a fully defined sequence of 192 amino acids amino acids as given in the specification. (15 pages)

L37 ANSWER 36 OF 105 HCAPLUS COPYRIGHT 2010 ACS on STN

AB Modified apoaequorin polypeptides that exhibit enhanced glowing are provided, as are functional fragments of the polypeptides, and polynucleotides encoding the polypeptides and functional fragments. A modified glowing aequorin photoprotein also is provided. In addition, methods of using the glowing apoaequorins and encoding polypeptides to detect, for example, the presence of calcium ions in a sample, or to identify agents that effect the movement of calcium ions from one compartment to another are provided, as are methods for identifying functional changes in cells associated with changes in calcium ion concns. Claimed cDNA sequences for aequorin were missing at time of publication.

L37 ANSWER 43 OF 105 MEDLINE on STN DUPLICATE 26

AB The Ca(2+)-regulated photoprotein obelin was substituted at Trp92 by His, Lys, Glu, and Arg. All mutants fold into stable conformations and produce bimodal bioluminescence spectra with enhanced contribution from a violet emission. The W92R mutant has an almost monomodal bioluminescence ($\lambda_{\text{max}}=390 \text{ nm}$) and monomodal fluorescence ($\lambda_{\text{max}}=425 \text{ nm}$) of the product. Results are interpreted by an excited state proton transfer mechanism involving the substituent side group and His22 in the binding cavity.

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AB A three-dimensional structure model was proposed for obelin, a photoprotein from hydroid polyp *Obelia longissima*. Alignment was applied to transform the amino acid sequence of invertebrate sarcoplasmic Ca²⁺-binding protein into the sequence of obelin, and the energy of the resulting protein was minimized. The model protein is a compact globule with a hydrophobic core. It has a cavity lined with residues important for photoprotein activity. The volume of the cavity is sufficient for binding the cofactor, implying that it contains the active center of the photoprotein. Several amino acid residues of obelin were

selected for mutational analysis.

L37 ANSWER 88 OF 105 HCPLUS COPYRIGHT 2010 ACS on STN
AB A model of the 3D structure of obelin-a photoprotein from sea organism *Obelia longissima* is suggested. Based on a 3D profile search of the compatibility of the sequence of photoproteins with the structure of calmodulin, troponin C, parvalbumin, and sarcoplasmic calcium binding protein (SCBP), the latter was chosen as template for modeling of the 3D structure of photoproteins. After substitution of the amino acid sequence of SCBP to that of obelin according to the alignment of their primary and secondary structures, the model was subjected to some rounds of energy minimization and the model obtained was analyzed. The structure contains a cavity which is lined by residues that have been shown to be important for the bioluminescence of photoproteins. To prove the suggested 3D structure of photoproteins and the suggested binding site for the photoactive compound some residues are proposed for mutational expts.

L37 ANSWER 95 OF 105 MEDLINE on STN DUPLICATE 50
AB Modification studies of the 5 histidine residues in aequorin employing site-directed mutagenesis and diethyl pyrocarbonate suggested that His169 may be the site of binding of molecular oxygen in aequorin. The modification of this residue led to complete loss of activity, whereas modification of the remaining 4 histidine residues yielded mutant aequorins with varying bioluminescence activities.

L37 ANSWER 105 OF 105 BIOSIS COPYRIGHT (c) 2010 The Thomson Corporation on STN

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L42 ANSWER 1 OF 4 WPIDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI Identifying substances that modulate nitric oxide release comprises contacting a test substance with a mixture of nitric oxide producer cells and detector cells having a reporter gene construct for detecting nitric oxide
PI DE 102006038942 A1 20080221 (200818)* DE 17[6]
WO 2008019783 A1 20080221 (200818) DE --
RW: AT BE BG BW CH CY CZ DE DK EA EE ES FI FR GB GH GM GR HU IE IS IT
KE LS LT LU LV MC MT MW MZ NA NL OA PL PT RO SD SE SI SK SL SZ TR
TZ UG ZM ZW
W: AE AG AL AM AT AU AZ BA BB BG BH BR BW BY BZ CA CH CN CO CR CU CZ
DE DK DM DO DZ EC EE EG ES FI GB GD GE GH GM GT HN HR HU ID IL IN
IS JP KE KG KM KN KP KR KZ LA LC LK LR LS LT LU LY MA MD ME MG MK
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SK SL SM SV SY TJ TM TN TR TT TZ UA UG US UZ VC VN ZA ZM ZW
IN STRAYLE J

L42 ANSWER 2 OF 4 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI New stable recombinant stem cell that expresses an apophotoprotein and produces a bioluminescent signal in the presence of a suitable chromophore, useful for in vitro testing of toxicity and/or teratology of a substance;
involving transgenic animal production expressing a apophotoprotein,
useful for a drug screening application
AU CAINARCA S; NUCCI C; CORAZZA S; LOHMER S
AN 2007-18390 BIOTECHDS
PI WO 2007080622 19 Jul 2007

L42 ANSWER 3 OF 4 BIOTECHDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI New non-human transgenic animal, useful for bio-imaging studies, for crossing with pharmacological relevant animal models, or for in vivo testing of toxicity and/or teratology of a substance;
involving trangenic mouse construction, useful for a transplantation experiment and as an animal model for a drug screening application
AU NUCCI C; CORAZZA S; LOHMER S
AN 2007-18389 BIOTECHDS
PI WO 2007080621 19 Jul 2007

L42 ANSWER 4 OF 4 WPIDS COPYRIGHT 2010 THOMSON REUTERS on STN
TI New nucleic acid encoding variant of aequorin, useful e.g. as reporter gene and as dye, has much longer luminescent lifetime than the parent protein, also new encoded proteins
PI DE 102005022146 A1 20061123 (200701)* DE 22[5]
WO 2006122650 A2 20061123 (200701) DE --
RW: AT BE BG BW CH CY CZ DE DK EA EE ES FI FR GB GH GM GR HU IE IS IT
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UG ZM ZW
W: AE AG AL AM AT AU AZ BA BB BG BR BW BY BZ CA CH CN CO CR CU CZ DE
DK DM DZ EC EE EG ES FI GB GD GE GH GM HR HU ID IL IN IS JP KE KG
KM KN KP KR KZ LC LK LR LS LT LU LV LY MA MD MG MK MN MW MX MZ NA
NG NI NO NZ OM PG PH PL PT RO RU SC SD SE SG SK SL SM SY TJ TM TN
TR TT TZ UA UG US UZ VC VN YU ZA ZM ZW
EP 1881992 A2 20080130 (200810) DE
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MC NL PL PT RO SE SI SK TR

IN 2007DN08638 P1 20071214 (200818) EN
CN 101223188 A 20080716 (200858) ZH
CA 2608004 A1 20061123 (200864) EN
KR 2008021018 A 20080306 (200864) KO
JP 2008539741 W 20081120 (200879) JA 33
TW 2007016177 A 20070501 (200934) ZH
US 20090203888 A1 20090813 (200954) # EN
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